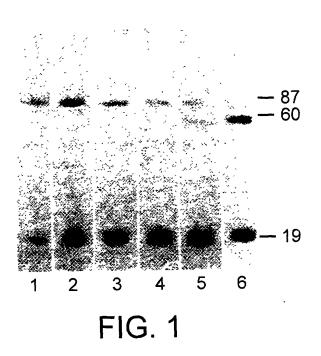
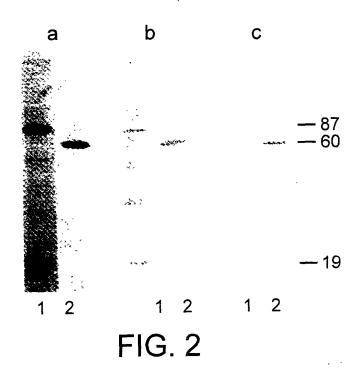
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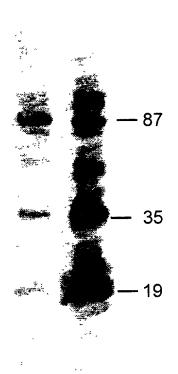


FIG. 3

2

1

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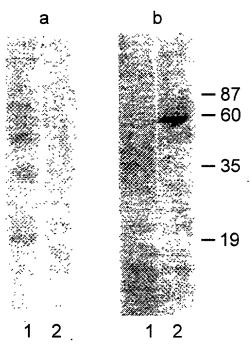
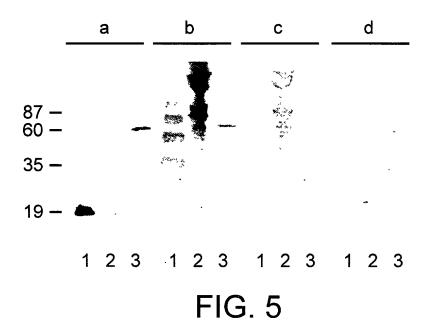


FIG. 4

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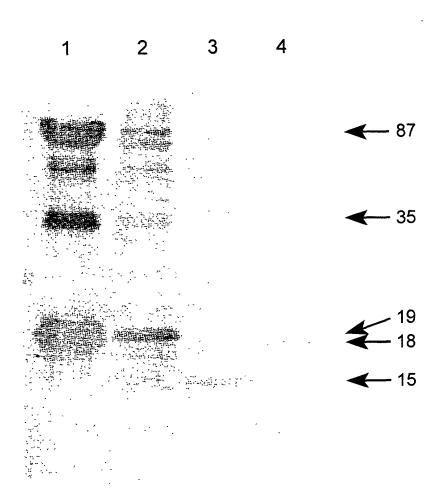
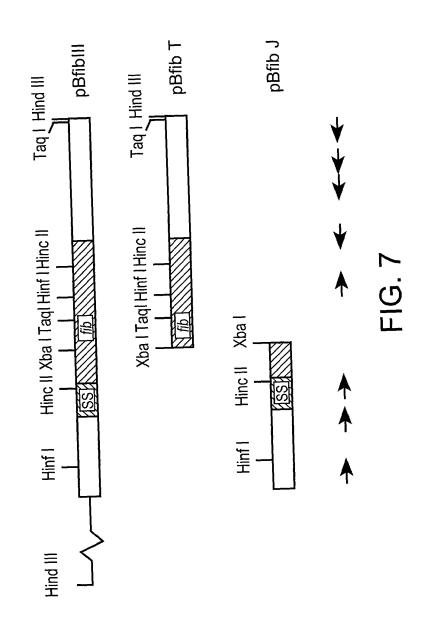


FIG. 6

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1 GACTAGTGTATAAGTGCTGATGAGTCACAAGATAGATAACTATATTTTGTCTATATTATA 60 -35 61 AAGTGTTTATAGTTAATTAATTAG<u>TTA</u>ATTTCAAAAGT<u>TG</u>TATAAATAGGATAACTT 120 -18 -35 121 AATA<u>AATGTA</u>AGATAATATTTGGAGGATAATTAACATGAAAAATAAATTGATAGCAAAA 180 -18 MKNKLIAK 181 TCTTTATTAACAATAGCGGCAATTGGTATTACTACAACTACAATTGCGTCAACAGCAGAT 240 SLLTIAAIGITTTTIASTAD 241 GCGAGCGAAGGATACGGTCCAAGAGAAAAGAAACCAGTGAGTATTAATCACAATATCGTA 300 A[†]SEGYGPREKKPVSINHNI 301 GAGTACAATGATGGTACTTTTAAATATCAATCTAGACCAAAATTTAACTCAACACCTAAA 360 EYNDGTFKYQSRPKFNSTPK 361 TATATTAAATTCAAACATGACTATAATATTTTAGAATTTAACGATGGTACATTCGAATAT 420 YIKFKHDYNILEFNDGTFEY 421 GGTGCACGTCCACAATTTAATAAACCAGCAGCGAAAACTGATGCAACTATTAAAAAAAGAA 480 GARPQFNKPAAKTDATIKKE 481 CAAAAATTGATTCAAGCTCAAAATCTTGTGAGAGAATTTGAAAAAACACATACTGTCAGT 540 QKLIQAQNLVREFEKTHTVS 541 GCACACAGAAAGCACAAAAGGCAGTCAACTTAGTTTCGTTTGAATACAAAGTGAAGAAA 600 AHRKAQKAVNLVSFEYKVKK 601 ATGGTCTTACAAGAGCGAATTGATAATGTATTAAAACAAGGATTAGTGAGATAATACTTC 660 M V L Q E R I D N V L K Q G L V R * 721 TATGCTATTTAATGCCAGGTTAATGTAACTTTCCTAAAATTGACTATATAATCGTTAAGT 780 MKFKKYILTGTLA 841 CATTACTTTTATCATCAACTGGGATAGCAACTATAGAAGGGAATAAAGCAGATGCAAGTA 900 LLLSSTGIATIEGNKADAS 901 GTCTGGACAAATATTTAACTGAAAGTCAGTTTCATGATAAACGCATAGCAGAAGAATTAA 960 LDKYLTESQFHDKRIAEELR 961 GAACTTTACTTAACAAATCGAATGTATATGCATTAGCTGCAGGAAGCTT 1009 TLLNKSNVYALAAGS

FIG. 8

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1	G CA	60 60
61 61	TTTCAAAAGTTGTATAAATAGGATAACTTAATAAATGTAAGATAATATTTGGAGGATAA	120 120
121 121	TTAACATGAAAAATAAATTGATAGCAAAATCTTTATTAACAATAGCGGCAATTGGTATTA G GC T A	180 180
181 181	CTACAACTACAATTGCGTCAACAGCAGATGCGAGCGAAGGATACGGTCCAAGAGAAAAGA	240 240
241 241	AACCAGTGAGTATTAATCACAATATCGTAGAGTACAATGATGGTACTTTTAAATATCAAT	300 300
301 301	CTAGACCAAAATTTAACTCAACACCTAAATATATTAAATTCAAACATGACTATAATATTT	360 360
361 361	TAGAATTTAACGATGGTACATTCGAATATGGTGCACGTCCACAATTTAATAAACCAGCAG	420 420
421 421	CGAAAACTGATGCAACTATTAAAAAAGAACAAAATTGATTCAAGCTCAAAATCTTGTGA	480 480
481 481	GAGAATTTGAAAAAACACATACTGTCAGTGCACACAGAAAAGCACAAAAGGCAGTCAACT	540 540
541 541	TAGTTTCGTTTGAATACAAAGTGAAGAAAATGGTCTTACAAGAGCGAATTGATAATGTAT	600 600
601 601	TAAAACAAGGATTAGTGAGATAATACTTCTGTCATTATTTTAAGTTCAAAATAATT T A A AA C GC G T TC GG TAAT	660 660
661 661	TAATATTATTATTTATTAATAAAACGACTATGCTATTTAATGCCAGGTTAATGTA A G G A G A G AT A	720 720
721 721	ACTTTCCTAAAATTGACTATAATCGTTAAGTATCAATTTTAAGGAGAGTTTACAATGA T G G C T	780 780
781 781	AATTT 785 785	

FIG. 9

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1 1	MKNKLIAKSLLTIAAIGITTTTIASTADASEGYGPREKKPVSINHNIVEYNDGTFKYQSR A L	60 60
61 61	PKFNSTPKYIKFKHDYNILEFNDGTFEYGARPQFNKPAAKTDATIKKEQKLIQAQNLVRE	120 120
121 121	FEKTHTVSAHRKAQKAVNLVSFEYKVKKMVLQERIDNVLKQGLVR 165 K 165	

FIG. 10

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Fbg-bp, strain Newman:

SEGYGPR

EKKPVSINH

YOSRPKFNSTP YGARPQFNKPA NIVEYNDGSFK NILEFINDGEFE KYIKFKHDY AKTDATIKKEQKLIQAQNLVREFEKTHTVSAHRKAQKAVNLVSFEYKVKKMVLQERIDNVLKQGLVR

Coagulase, strain 8325-4:

(C-terminal fragment)

YGPRPQFNKTP ASQ

YGARPTYKKPS YEARPRENKPS GIREYNDGTFG NVTTHANGOVS

ETNAY

KYVKYRDAGT

KTNAY

ETNAY

KTNAY KTNAY

YGARPTONKPS YGARQAQNKPS NVTTHANGOVS NVTTHGNGOVS

YGARPTYKKPS YGPRVTK NVTTHADGEAT NVTTHANGQVS

FIG. 11

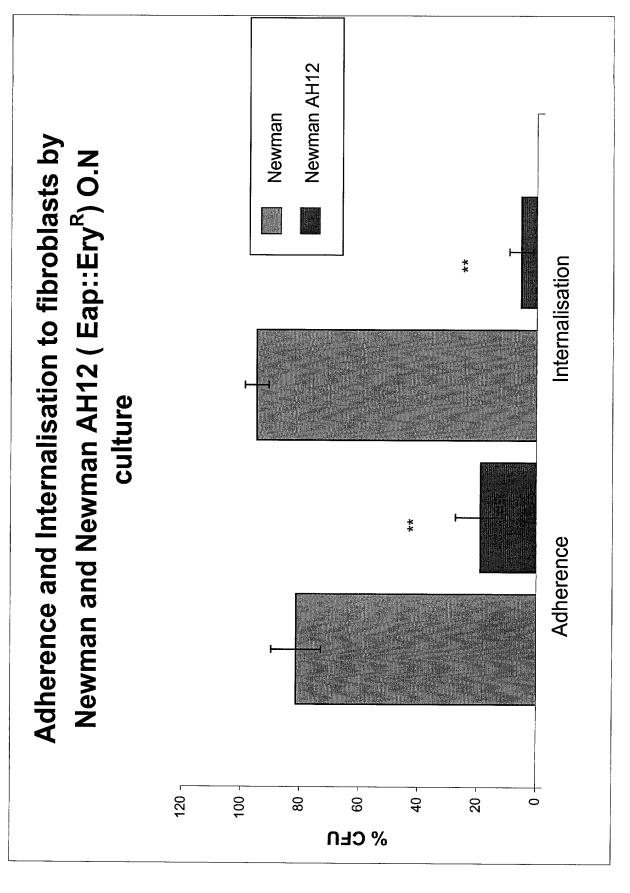


FIG. 12

FIG. 13

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FIG. 14

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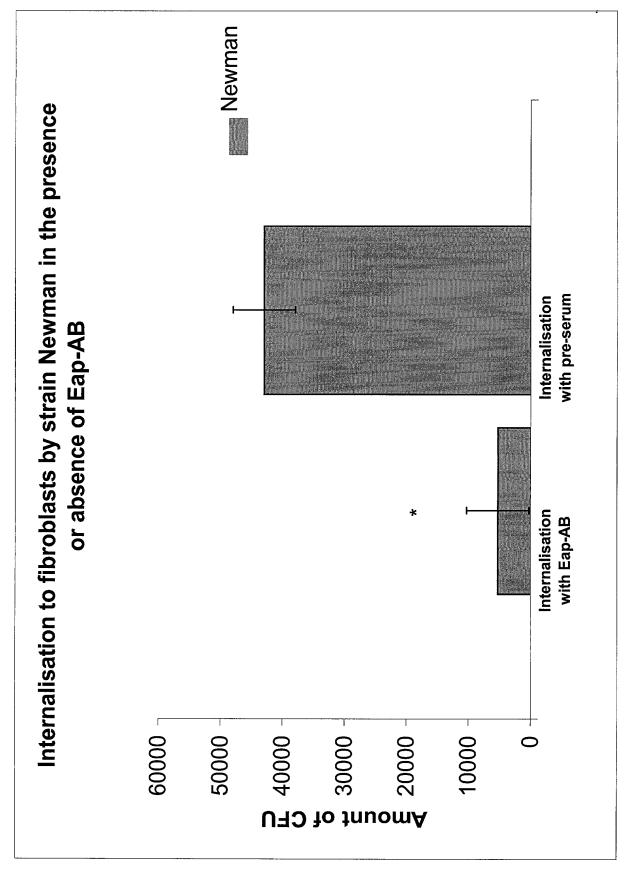


FIG. 15